

Figure 1

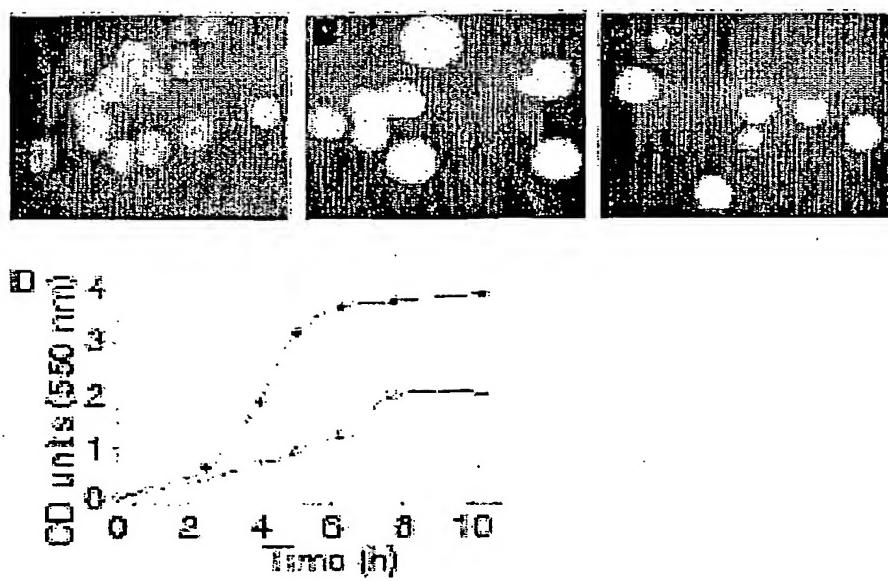


Figure 2

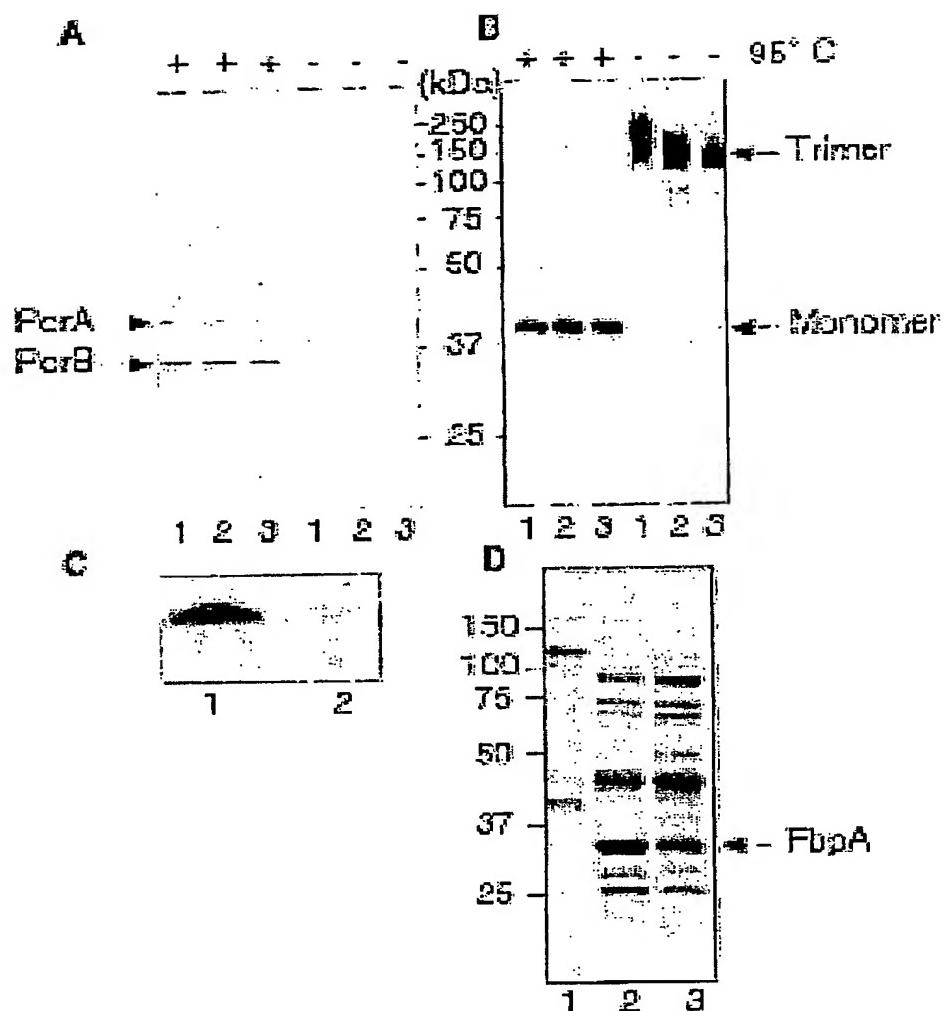


Figure 3

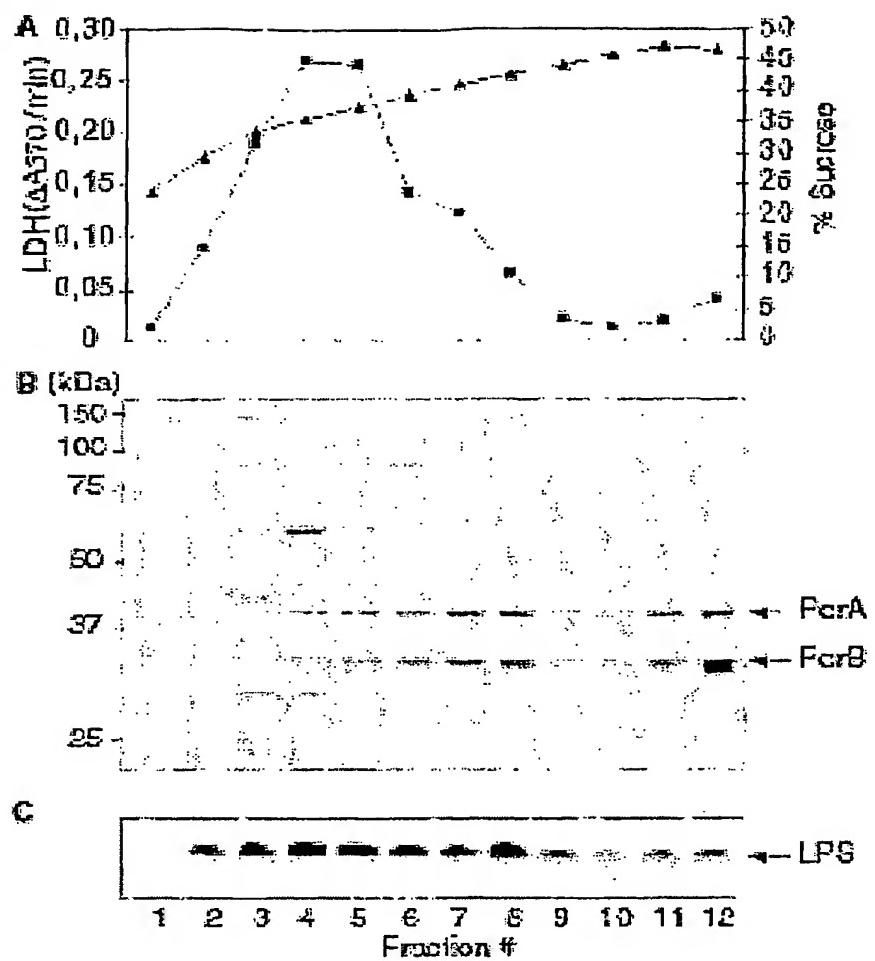


Figure 4

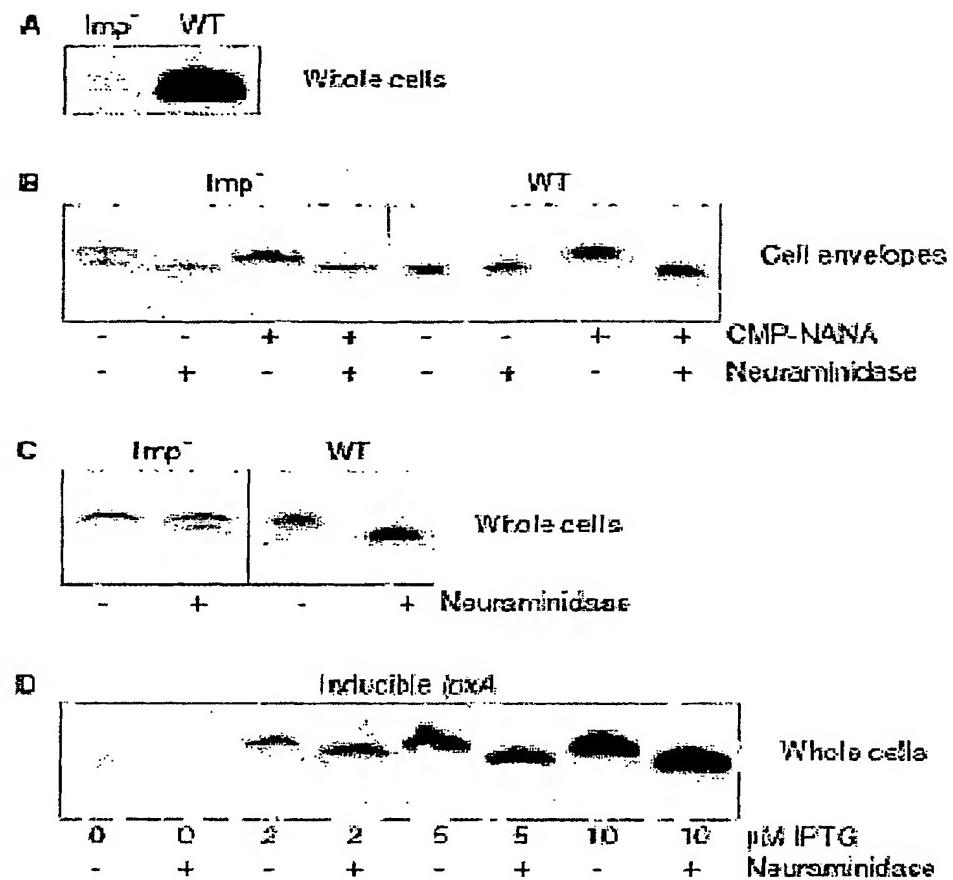


Figure 5

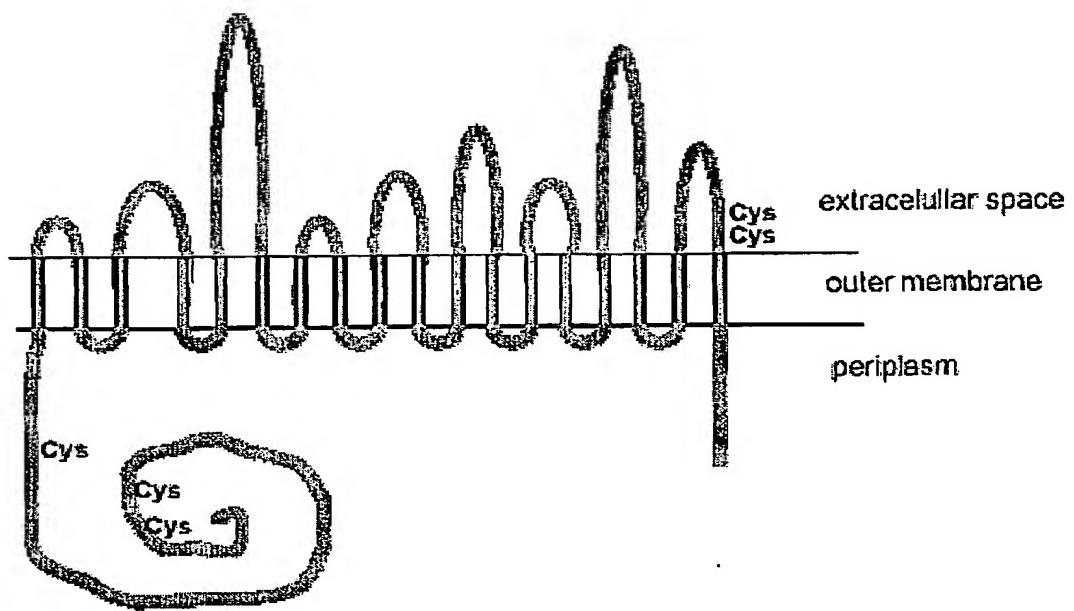


Figure 6

Imp/OstA - SEQ ID No 1

Predicted extracellular loops are indicated in bold

LARLFSLKPLVLALGLCFGTHCAAADAVAAEETDNPTAGESVRSVSEPIQPTSLSLGSTC 60
LFCNSESGSPERTEAAVQGSGEASIPEDYTRIVADRMEGQSQVQVRAEGNVVVERNRTTL 120
NTDWADYDQSGDTVTAGDRFALQQDGTЛИRGETLTYNLEQQTGEAHNVRMEIEQGGRRLQ 180
SVSRTAEMLGEGHYKLTETQFNTCSAGDAGWYVKAASVEADREKGIGVAKHAAVFVFGGP 240
IFYTPWADFPLDGNRKSGLLVPSLSAGSDGVSLSPYYFNLAPNLDATFAPSVIGERGAV 300
FDGQVRYLRPDYAGQSDLTWLPHDKSGRNNRYQAKWQHRHDISDTLQAGVDFNQVSDSG 360
YYRDFYGNKEIAGNVNLNRRVWLGYGRAAGGSLNAGLSVLKYQTLANQSGYKDCKPYALM 420
PRLSVEWRKNTGRAQIGVSAQFTRFSHDSRQDGSRLVYPDIKWDFSNWGYVRPKLGLH 480
ATYYSLNRFGSQEARRVSRTLPIVNIDSGATFERNTRMFGGEVLQTLPEPLFYNYIPAKS 540
QNDLPNFDSSESSFYGYGQLFRENLYYGNDRINTANSLSAAVQSRILDGATGEERFRAGIG 600
QKFYFKDDAVMLGSVGKKPRNRSDWVAFASGSIGSRFILDSSIHYNQNDKRAENYAVGA 660
SYRPAQGKVNLARYKYGRNEKIYLKSDGSYFYDKLSQDLSAQWPLTRNLSAVVRYNYGF 720
EAKKPIEVLAGAEYKSSCGCWGAGVYAQRYVTGENTYKNAVFFSLQLKDLSSVGRNPADR 780
MDVAVPGYITAHSLAGRNKRP 802

loop 1: 252-271
loop 2: 295-332
loop 3: 357-416
loop 4: 444-455
loop 5: 482-501
loop 6: 537-576
loop 7: 606-624
loop 8: 648-697
loop 9: 721-740

Figure 7

Imp - FAM18 NMB0280 NMA2207	LARLFLSLKPLVLALGFCFGTHCAAADAVAAEETDNPTAGGSVRSVSEPIQPTSLSLGSTC LARLFLSLKPLVLALGFCFGTHCAAADAVAAEETDNPTAGESVRSVSEPIQPTSLSLGSTC MARLFSLKPLVLALGFCFGTHCAAADAVAAEETDNPTAGGSVRSVSEPIQPTSLSLGSTC *****	60 60 60 *****
Imp - FAM18 NMB0280 NMA2207	LFCSNESGGSPERTEAAVRGSGEASI PEDYTRIVADKVEGQSQVQVRAEGNVVVERNRTTL LFCSNESGGSPERTEAAVQGSGEASI PEDYTRIVADRMEGQSQVQVRAEGNVVVERNRTTL LFCSNESGGSPERTEAAVQGSGEASI PEDYTRIVADRMEGQSQVQVRAEGNVVVERNRTTL *****	120 120 120 *****
Imp - FAM18 NMB0280 NMA2207	NTDWADYDQSGDTVTAGDRFALQDGTLIRGETLTYNLEQQTGEAHNVRMETEHGGRRLQ NTDWADYDQSGDTVTAGDRFALQDGTLIRGETLTYNLEQQTGEAHNVRMEIEQGGRRLQ NADWADYDQSGDTVTAGDRFALQDGTLIRGETLTYNLEQQTGEAHNVRMETEHGGRRLQ *****	180 180 180 *****
Imp - FAM18 NMB0280 NMA2207	SVSRTAEMLGEGHYKLTELQFNTCSAGDAGWVKAASVEADREKGIGVAKHAAFVFGGP SVSRTAEMLGEGHYKLTELQFNTCSAGDAGWVKAASVEADREKGIGVAKHAAFVFGGP SVSRTAEMLGEGHYKLTELQFNTCSAGDAGWVKAASVEADREKGIGVAKHAAFVFGGP *****	240 240 240 *****
Imp - FAM18 NMB0280 NMA2207	IFYTPWADFPLDGNRKSGLLVPSLSAGSDGVSLSPVYYFLAPNL DATFAPSVIGERAV IFYTPWADFPLDGNRKSGLLVPSLSAGSDGVSLSPVYYFLAPNL DATFAPSVIGERAV IFYTPWADFPLDGNRKSGLLVPSLSAGSDGVSLSPVYYFLAPNL DATFAPGVIGERAV *****	300 300 300 *****
Imp - FAM18 NMB0280 NMA2207	FDGQVRYLRPDYAGQSDLTLPHDKKSGRNMRYQAKWQHRDISDTLQAGVDFNQVSDSG FDGQVRYLRPDYAGQSDLTLPHDKKSGRNMRYQAKWQHRDISDTLQAGVDFNQVSDSG FDGQVRYLRPDYAGQSDLTLPHDKKSGRNMRYQAKWQHRDISDTLQAGVDFNQVSDSG *****	360 360 360 *****
Imp - FAM18 NMB0280 NMA2207	YYRDFYGNKEIAGNVNLNRVLDYGRAAGGSLNAGLSVLKYQTIANQSGYKDCKYALM YYRDFYGNKEIAGNVNLNRVLDYGRAAGGSLNAGLSVLKYQTIANQSGYKDCKYALM YYRDFYGNKEIAGNVNLNRVLDYGRAAGGSLNAGLSVLKYQTIANQSGYKDCKYALM *****	420 420 420 *****
Imp - FAM18 NMB0280 NMA2207	PRLSADWRKNTGRAQIGVSAQFTRFSHDSRQDGSLRVVYDPIKWDFTNSNWGYVRPKLGLH PRLSVEWRKNTGRAQIGVSAQFTRFSHDSRQDGSLRVVYDPIKWDFTNSNWGYVRPKLGLH PRLSADWRKNTGRAQIGVSAQFTRFSHDSRQDGSLRVVYDPIKWDFTNSNWGYVRPKLGLH *****	480 480 480 *****
Imp - FAM18 NMB0280 NMA2207	ATYYSLNRFGSQEARRSRTLPIVNIDSGATFERNTRMFGGVLQITLEPRLFYNYIPAKS ATYYSLNRFGSQEARRSRTLPIVNIDSGATFERNTRMFGEVVLQITLEPRLFYNYIPAKS ATYYSLNRFGSQEARRSRTLPIVNIDSGMTFERNTRMFGGVLQITLEPRLFYNYIPAKS *****	540 540 540 *****
Imp - FAM18 NMB0280 NMA2207	QNDLPNFDSSESSFYQQLFRENLYYGNDRINTANSLSAAVQSRILDGATGEERFRAGIG QNDLPNFDSSESSFYQQLFRENLYYGNDRINTANSLSAAVQSRILDGATGEERFRAGIG QNDLPNFDSSESSFYQQLFRENLYYGNDRINTANSLSAAVQSRILDGATGEERFRAGIG *****	600 600 600 *****
Imp - FAM18 NMB0280 NMA2207	QKFYFKDDAVMLDGSVGKKPRNRSRWVAFASGSIGSRFIELDSSIHYNQNDKRAENYAVGA QKFYFKDDAVMLDGSVGKKPRNRSRWVAFASGSIGSRFIELDSSIHYNQNDKRAENYAVGA QKFYFKDDAVMLDGSVGKKPRRSRWVAFASSGIGSRFIELDSSIHYNQNDKRAENYAVGA *****	660 660 660 *****
Imp - FAM18 NMB0280 NMA2207	SYRPAQGVNLNARYKGRNEKIYLKSDGSFYDKLSQLDLSAQWPTRNLSAVVRYNYGF SYRPAQGVNLNARYKGRNEKIYLKSDGSFYDKLSQLDLSAQWPTRNLSAVVRYNYGF SYRPAQGVNLNARYKGRNEKIYLKSDGSFYDKLSQLDLSAQWPTRNLSAVVRYNYGF *****	720 720 720 *****
Imp - FAM18 NMB0280 NMA2207	EAKKPIEMLAGAEYKSSCGCWGAGVYAQRYVTGENTYKNAVFFSLQLKDLSSVGRNPADR EAKKPIEVLAGAEYKSSCGCWGAGVYAQRYVTGENTYKNAVFFSLQLKDLSSVGRNPADR EAKKPIEVLAGAEYKSSCGCWGAGVYAQRYVTGENTYKNAVFFSLQLKDLSSVGRNPADR *****	780 780 780 *****
Imp - FAM18 NMB0280 NMA2207	MDVAVPGYIPAHSLSAGRNP 802 MDVAVPGYITAHSLSAGRNP 802 MDVAVPGYIPAHSLSAGRNP 802 *****	802 802 802 *****

Figure 8

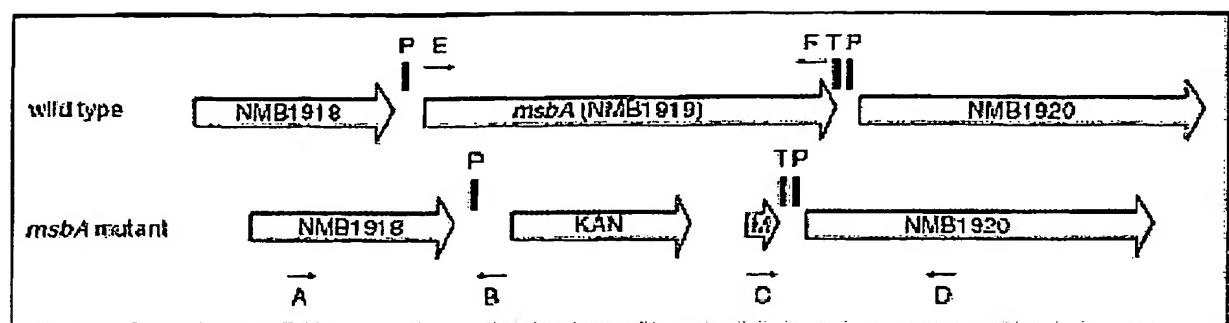


Figure 9

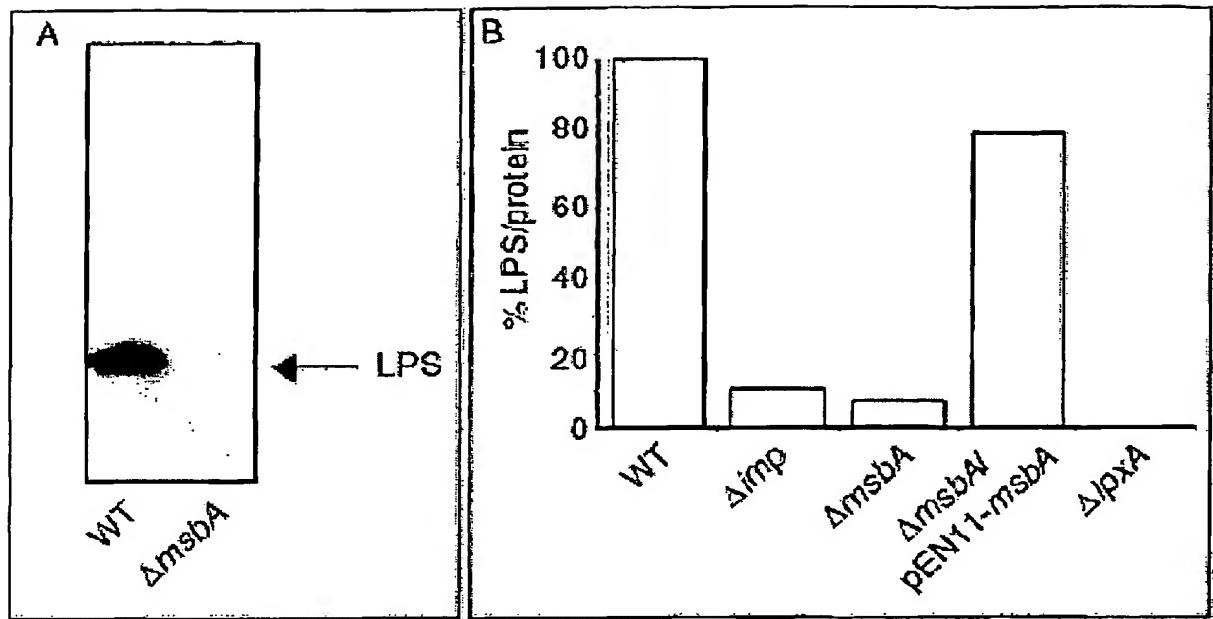


Figure 10

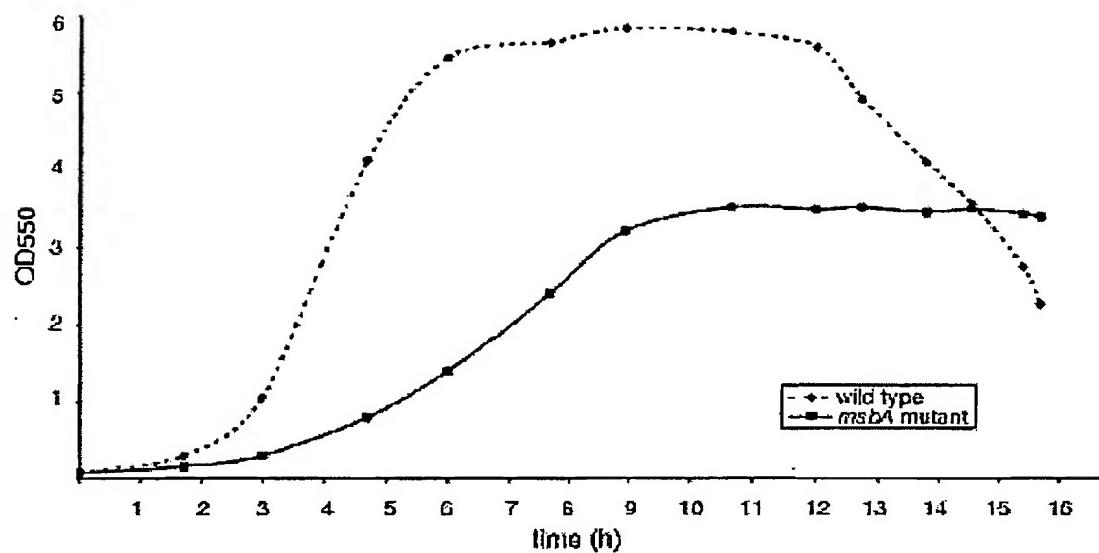


Figure 11

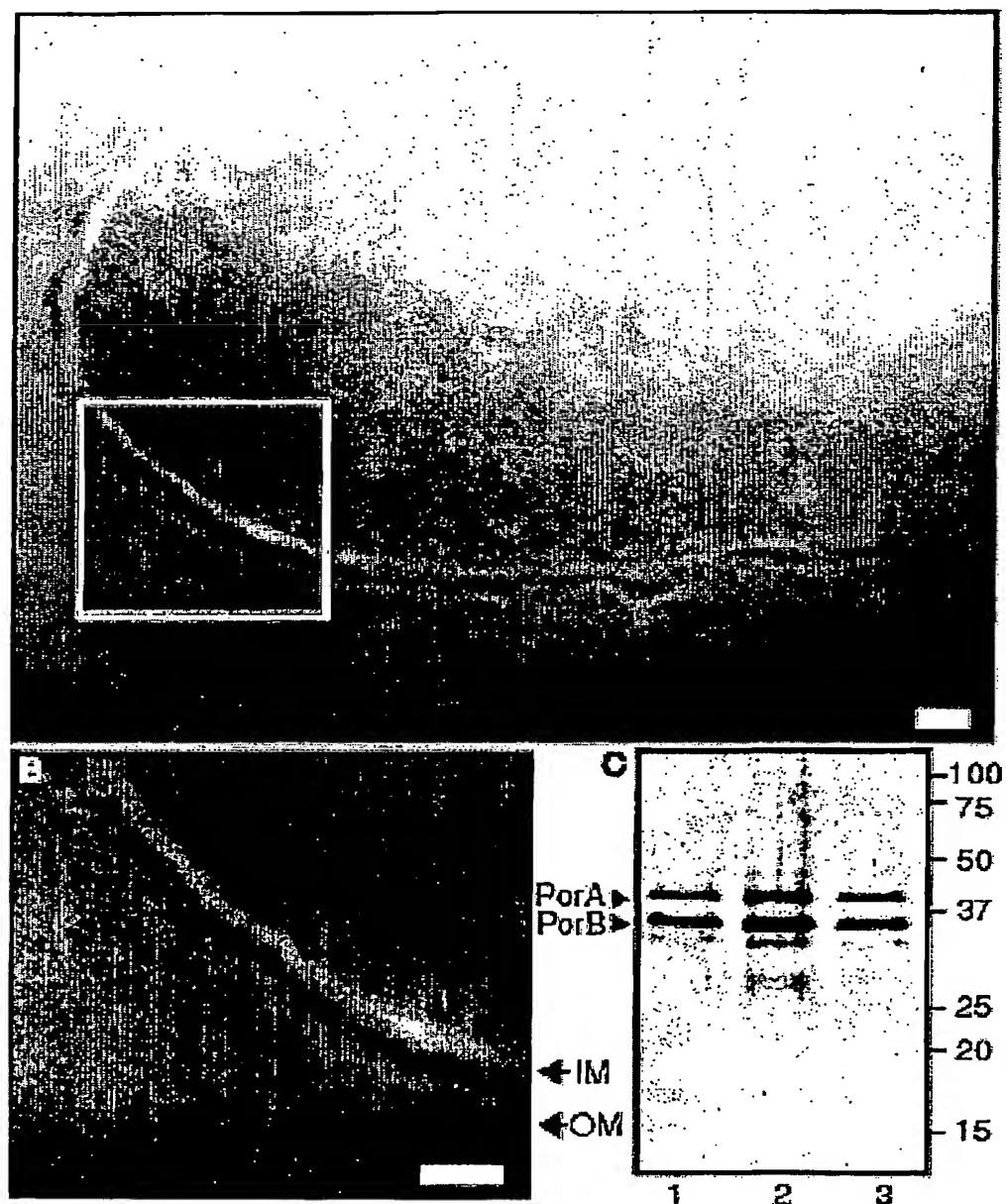


Figure 12

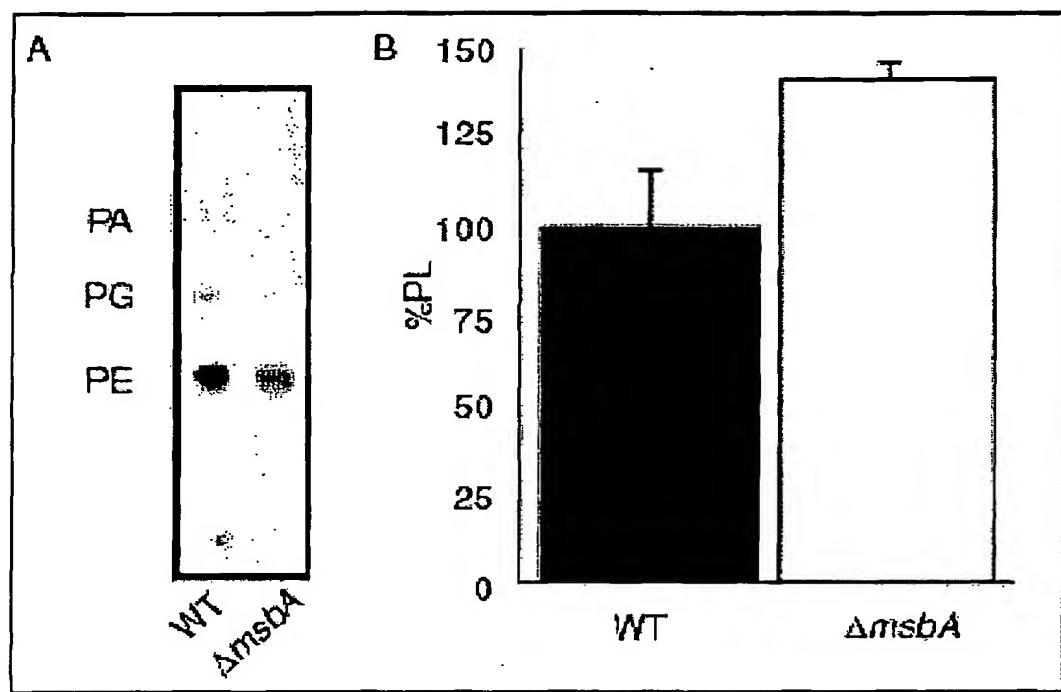


Figure 13

A

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B

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Figure 14

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Figure 15 A

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Figure 15 B